Figure \_\_

Nucleotide and Amino Acid Sequence of the h3D1 Light Chain

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TCTAGACCACC ATG GAT TCA CAG GCC CAG GTT CTT ATA
Xbal
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 рl
                          TCT GGC ACC
                                       TGT
                                           GGG GAC
                                                    ATT
                                                        GIG
          CTG CTA TGG GTA
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                                    T
                                             G
                                           GTA AGC
                                                        GGA
              CAG TCT
                      CCA
                          GAT TCC CTG
                                       GCT
                                                    TTA
         ACA
      CTG
p24 ▶
               Q
                                                     L
                          AGC
                               TGC AAA
                                       TCC
                                            AGT CAG
              GCC ACT ATT
         AGG
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                                                 Q
p37 •
                       I
                                C
           R
                                                    TCC
                                                GCT
                                                         TAC
      CTC AAC
              AGT AGA ACC
                           CGA
                               GAG
                                   AAC
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                                                CLC
                                                    ATC
      CAG CAG
              AAA CCA GGG
                           CAG CCT
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p63 ▶ Q
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                       G
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                                        GTC
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                                                         TIC
                           GAA TCT GGG
      TGG GCA
              TCC ACT AGG
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p76 🕨
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                           GGG ACA GAT
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                                                CIC ACC
                                                         ATC
      AGT GGC AGT GGA TCT
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p89 🕨 S
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                                       GCA GTT
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      AGC AGT
              CTG CAG GCT GAA GAC GTG
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          CAA TCT TAT AAT CTT TAC ACG
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      AAG GTG GAA ATA AAA C gtaagtagtetteteaactetagaaattetaa
p128 ▶ 🕱
                                                  Xpal
           V
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      actotgagggggtoggatgaogtggocattotttgcotaaagcattgagtttac
      tgcaaggtcagaaaagcatgcaaagccctcagaatggctgcaaagagctccaac
      aaaacaatttagaactttattaaggaatagggggaagctaggaagaaactcaaa
      acatcaagattttaaatacgcttcttggtctccttgctataattatctgggata
      agcatgctgttttctgtctgtccctaacatgccctgtgattatccgcaaacaac
      acacccaagggcagaactttgttacttaaacaccatcctgtttgcttctttcct
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(continued on the next page)

Figure Chain Gene (cont'd): Nucleotide and Amino Acid Sequence of the h3D1 Light

GA ACT GTG GCT GCA CCA TCT GTC TTC ATC P I p133 Þ S GGA ACT GCC TCT GTT CCA TCT GAT GAG CAG TTG AAA TCT p145 T v D ĸ 8 E Q L TAT CCC AGA GAG GCC AAA TGC CTG CTG AAT AAC TTC P E A p158 ¥ R C L L N N P CAG TGG AAG GTG GAT AAC GCC CTC CAA TCG GGT AAC GTA G v N L Q N p171 Q K GAC AGC AAG GAC AGC CAG GAG AGT GTC ACA GAG CAG TCC E 3 V E D p184 Q T 0 TAC AGC CTC AGC AGC ACC CTG ACG CTG AGC AAA GCA p197 Y S L 3 3 Ŧ L T L S ĸ λ TGC GAA GTC ACC GAC TAC GAG AAA CAC AAA GIC TAC GCC p210 🕨 ¥ C 15 v Y E ĸ Ħ ĸ v CAT CAG GGC CTG AGC TCG CCC GTC ACA AAG AGC TTC AAC p223 ₽ v T ĸ 2 AGG GGA GAG TGT TAG agggagaagtgcccccacctgctcctcagttc p236 > R E C Stop

cagcotgaccccctcccatcctttggcctctgaccctttttccacaggggac ctacccctattgcggtcctccagctcatctttcacctcacccccctcctcct ccttggctttaattatgctaatgttggaggagaatgaataaattaaagtgaa tctttgcacctgtggtttctctctttcctcatttaataattattatctgttg ttttaccaactactcaatttctcttataagggactaaatatgtagtcatcct aaggcgcataaccatttataaaaatcatccttcattctattttaccctatca toctotgcaagacagtcctccctcaaacccacaagccttctgtcctcacagt cccctgggccatggtaggagacttgcttccttgttttcccctcctcagca ageceteatagteetttttaagggtgacaggtettacagteatatateettt gattcaattccctgggaatcaaccaaagcaaatttttcaaaagaagaaacct gctataaagagaatcattcattgcaacatgatataaaataacaacacaataa aagcaattaaataaacaaacaatagggaaatgtttaagttcatcatggtact tagacttaatggaatgtcatgccttatttacatttttaaacaggtactgagg gactectgtetgeeaagggeegtattgagtaettteeacaacctaatttaat: ccacactatactgtgagattaaaaacattcattaaaatgttgcaaaggttct ataaagotgagagacaaatatattotataactcagcaatcccacttotagga tcaattc

CK

Figure Gene

## Nucleotide and Amino Acid Sequence of the h3D1 Heavy Chain

Xbal		מיוש אני	GACC	ACC	ATG	GGT	TGG	AAC	TGT	ATC	ATC	TTC	TT	r	TG	1	
•		47-44		•	M	G	W	N	c	I	ľ	F	P		L	1	
p1		GTT	AÇC	מיטמ	दिया	ACA	CCT	GIG	CAC	TCC	CAG	GIC	CA	G (	TG		
pll		A.	T	T	A	T	G	v	H	5	Q	v	Q		L	1	
p		GTG	CAG	TCT		GCT	GAG	GIG	AAG	AAG	CCT	GGG	AG	C f	ICA		
p24			Q	s	G	A	E	v	X	X	P	G	5		S		
P2-4	•	GIG.		GTG	TCC		AAA	GCT	TCC	GGC	TAC	ACA	TI	C:	ACT		
p37	Þ	T.	K	v	8	c	K	A	5	G	Y	T	F		T	- [	
ps,	•	GAT	••	GCT	ATA	CAG	TGG	GIG	AGA	CAG	GCT	CCT	GG	Α	CAG	Ţ	
p50	Þ		Y	A	I	Q	W	V	R	Q	A	P	G	<b>;</b>	Q	]	
•	٠	GGC	CIC	GAG	TGG	ATT	GGA	GTT	ATT	AAT	ATT	TAC	T	$\mathbf{T}$	GAT	- [ '	VH
p63	•	G	L	E	W	I	G	٧	I	N	I	Y	7	•	מ	- L	
Poo	•	AAT	ACA	AAC	TAC	AAC	CAG	AAG	TIT	AAG	GGC	AAG	GC	C	ACA	-1	
p76	•		T	N	Y	H	Q	K	F	ĸ	G	K	2	<b>L</b>	T	1	
•		ATG	ACT	GIA	GAC	AAC	TCG	ACG	AGC	ACA	GCC	TAT	A	ľG	GAA	- 1	
p89	•	M	T	v	D	K	S	T	S	T	A	Y	ŀ	1	E	1	
-		CTT	AGT	TCT	TIG	AGA	TCT	GAG	GAT	ACG				YΤ	TAC	1	
p102	Þ	L	S	8	L	R	s	E	D	T	A	٧		7	Y	1	
		<b>TGT</b>	GCA	AGA									_	JT.	CAA		
p115		C	y.	R	A	A	W	X	M	D	Y	W		3	Q		
400	_	CCI	-		CIC	ACC				G g	tgaç	iccc.	cta	aa		도 bai	
p128	,		T	L	v	T	V	5	s						^	Der I	
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		tag	agct	ttct	<b>gggg</b>	cgag	ccgg	gcct	gact	ttgg	ctt	ggg	gca	999	gagt stop	g	
		ggc	taag	gtga	ggca	ggtg	gcgc	cagc	cagg	tgca	CAC	caa	tgc	CC	gtga	g	
		ggc	taag	gtga actg	ggca gacc	ggtg ctgc	gcgc	cagc accc	cagg tcgt	tgca ggat	agac :agac	caa caag	tgc aac	cg	<b>z</b> ggg gtga	g	
		ggc	taag agac ctgc	gtga actg gccc	ggca gacc tggg	ggtg ctgc ccca	gcgc ctgg gctc	cagc accc tgtc	cagg tegt ¢cac	tgca ggat accg	icaal reacc	caa caag	tgc aac atg	g¢ cg	<b>z</b> ggg gtga	ig g	
n136		ggc	taag	gtga actg gccc	ggca gacc tggg	ggtg ctgc ccca	gcgc ctgg gctc	cagc accc	cagg tegt ¢cac	tgca ggat accg	icaal reacc	caa caag	tgc aac atg	g¢ cg	gtga 2ggg acca	ig g	
p136	ı	ggc ccc cct	taag agac ctgc tctt	gtga actg gccc gcag	ggca gacc tggg CC A	ggtg ctgc ccca TCC s	gege etgg gete ACC	cagc accc tgtc AAG	cagg tegt ccac GGC G	tgca ggat accg cca	LCACO LCGGI LCGGI LCGGI LCGGI LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO	caa aag cac GT V	tgc aac atg C I	cg gc TC	CCC acca aggg acca	ig G	
-		ggc ccc cct ctc	taag agac ctgc	gtga actg gccc gcag	ggca gacc tggg CC A	ggtg ctgc ccca TCC s	gege etgg gete ACC	cagc accc tgtc AAG	cagg tegt ccae GGC G	tgca ggat accg cca	LCACO LCGGI LCGGI LCGGI LCGGI LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO LCACO	caa aag cac G G V	tgc aac atg C I	cg gc TC	gtga acca CCC P	ig G	
p136 p146		ggc ccc cct ctc	taag agac ctgc tctt GCG	gtga actg gcag gcag CCC	ggca gacc tggg CC A TGC	ggtg ctgc ccca TCC s TCC	gege etgg gete ACC T ! AGG	cage accc tgtc AAG K AGC	cagg tegt ccac GGC G ACC	tgca ggat accg CCP TCC	LCACO LAGRA JCGGI A TCO S CAA LAGRA E	cead aag cac G GT V AG	tgc aac atg C I	cg gc TC F CA	CCC P CCC BGGG BGGG BGGG BGGG BGGG BGGG	g : :	
p146	)	CIG CCC CCC CCC CCC CCC	taag agac ctgc tctt GCG	gtga actg gcag gcag CCC	ggca gacc tggg CC A TGC	ggtg ctgc ccca TCC s TCC	gege etgg gete ACC T ! AGG	cagc accc tgtc AAG K AGC	cagg tegt ccac GGC G ACC	tgca ggat accg CCP TCC	LCACO LAGRA JCGGI A TCO S CAA LAGRA E	cead aag cac G GT V AG	tgc aac atg C T C A	cg gc TC F CA	CCC P CCC AGGG AGGG AGGG AGGG AGGG AGGG	g : :	
-	)	CIG CCC CCC CCC CCC CCC	taag agac ctgc tctt GCG A CTG	gtga actg gcag gcag GGC GGC	ggca tggg CC A TGC TGC	ggtg ccca TCC s TCC S CTC	gcgc gctc gctc ACC r AGG R GTC	cagc accc tgtc AAG K AGC 5 AAG	cagg tegt ccac GGC ACC T GAC	tgca ggat acco c CCP TCC S TAC	Leace Lagad Reggi L TCC S CAA E TTC	caa caag cac c or v c Ag s c cc	tgc aac atg C I	CG GG TC F LCA T	etga aggg acca CCC P GCC A CCC	g g c	
p146 p159	)	CTG CTG A GCC	taag agac ctgc tctt CCG A CTG	gtga actg gcag gcag GGC GGC	TGC TGC TGC TGC	ggtg ccca TCC s TCC S CTC	gcgc gctc gctc ACC r AGG R GTC	cagc accc tgtc AAG K AGC S AAG	cagg tegt ccac GGC ACC T GAC	tgca ggat acco c CCP TCC S TAC	Leace Lagad Reggi C GAA E TTV	caa caag cac c or v c Ag s c cc	tgc aac atg C T C A	cc gc TC F ACA T EAA	etga aggg acca CCC P GCC A CCC	g g c	СН1
p146	)	CTG CTG A GCC	taag agac ctgc tctt  GCG A CTG L ACG	gtga actg gccc gcag ccc P GGC GTG	ggca gacc tggg TGC C TCC	ggtg ctgc ccca TCC s TCC S CTG L	gege gete gete ACC T AGG R GTC V AAC	cage tgtc : AAG : AGC : AAG : AAG : TCA	cagg tegt ccae GGC ACC T GAC D GGC	tgca ggat acce ccr r TCC s TAC Y	LCACCACACACACACACACACACACACACACACACACAC	ceaa caag cac cor v cac cac cac cac cac cac cac cac cac c	tgc atg C T C A	CCP CCP CCP CCP CCP CCP CCP CCP CCP CCP	P CTT	9 6	СН1
p146 p159	1	GIG GIG GIG GIG GIG GIG GIG GIG GIG GIG	taag agac ctgc tctt  GCG A CTG L ACG	gtga actg gccc gcag gccc gcag yccc	ggca gacc tggg TGC C TCC	ggtg ctgc ccca TCC s TCC S CTG L	gege gete gete ACC T AGG R GTC V AAC	cage tgtc AAG AGO S AAG TCA	cagg tegt ccae GGC ACC T GAC D GGC	tgca ggat accg CCF TCC S TAC Y GCC A	LCACCACACACACACACACACACACACACACACACACAC	ceaa caag ceac coac y AG AG CC CC TC CT CT	tgc aac atg C T C A	CCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCC	E CTY	9 9 6 6	CH1
p146 p159 p172	1	GIG GIG GIG GIG GIG GIG GIG GIG GIG GIG	taag agac ctgc tctt GCG A CTG L ACG T	gtga actg gccc gcag gccc gcag GCC GCC GCC GCC GCC GCC GCC GCC GCC GC	ggca gacc tggg CC TGC C TCC s	ggtg ccca TCC S TCC S CTG L G TGG	gege gete gete ACC AGG R GTC V AAC N ACC A	cage tgtc : AAG : AGC : AAG : AAG : TCA : TCA	cagg tegt ccac GGC GAC T GAC D GGC GCIII	tgca ggat acco P TCC S TAX S CAA CAA	Leace Lagad Reggi Leace CAM CAM F CT T CT CT CT CT CT CT CT CT CT CT CT C	ccaa caag ccac c cac c c cac c c cac c c cac c c cac c c c c	tgc aac atg C T C A C A C A C A C A C A C A C A C A C A	CORRECTE STATE OF STA	E CTC  CTC  CTC  CTC  CTC  CTC  CTC  CTC	9 9 6 6	CH1
p146 p159 p172	) ! ! !	ggc ccc ctc CTG CTG CTG	taag agac ctgc tctt GCG A CTG L ACG T	gtga actg gccc gcag gccc gcag GCC GCC GCC GCC GCC GCC GCC GCC GCC GC	ggca gacc tggg CC TGC C TCC s	ggtg ccca TCC s TCC S CTG L GGG W	gege gete gete ACC AGG R GTC V AAC N ACC A	E GTO	Cagg coac GGC GCC ACC GAC GAC CTI L GAC T	tgca ggat acco P TOO S TAO Y GOT A A O A	Leace Leagan Leagan Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leach Leac	COAA CAAGGACAACAACAACAACAACAACAACAACAACAACAAC	tgc aac atg C T C C C C C C C C C C C C C C C C C C	CONTRACT SANGER	E COC COC A COC A COC COC COC COC		CH1
p146 p159 p172 p185 p198		ggc ccc ctc ctc CTG CTG GCC A GTG	taag agac ctgc tett GCG A CTG L ACG T CAC H	gtga actg gccc gcag gccc F GGC GGC T CTC T	ggca gacc tgggg CC A TGC TCC TCC S TTCC S TTCC	ggtg ctgc TCC S TCC S CTG L G TGC W	gege gete gete gete ACC R R GTC V AAC ACC N CCT V CTAC	E GTO ACC	Cagg coac GGC GCC ACC GAC GAC CTI L GAC T	tgca ggat acco cor TOC S TAC S TAC CAC CAC CAC CAC CAC CAC CAC CAC CAC	Leace cage cage cage cage cage cage cage	E AGE CO TO	tgc aacg C I C A C C A C C A C C C A C C C C C C	CAN	E CTO  COCO  COCO		СН1
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(continued on the next page)

## Figure Chain Gene

(cont'd): Nucleotide and Amino Acid Sequence of the h3D1 Heavy

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